## SEQUENCE LISTING

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<110> KUFER, PETER
     LENKKERI-SCHUTZ, ULLA
     LUTTERBUSE, RALF
     KOHLEISEN, BIRGIT
<120> LESS IMMUNOGENIC BINDING MOLECULES
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<140> 10/588,734
<141> 2006-08-08
<150> PCT/EP05/001573
<151> 2005-02-16
<150> EP 04003445.6
<151> 2004-02-16
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atcacttgca gtgcaagttc aagcgtaagc tacatgaatt ggtatcagca gacaccaggg
                                                                      120
aaagccccta agagatggat ctatgacaca tccaaattgg cttctggggt cccatcaagg
                                                                      180
ttcagtggca gtggatctgg gacagattac actttcacca tcagcagtct gcaacctgaa
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gatattgcaa cttactactg tcaacagtgg agtagtaacc cttttacttt tggccagggg
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accaagetge agateace
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Asp Arg Val Thr Ile Thr Cys Ser Ala Ser Ser Ser Val Ser Tyr Met
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25

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Asn Trp Tyr Gln Gln Thr Pro Gly Lys Ala Pro Lys Arg Trp Ile Tyr
                            40
                                                 45
        35
Asp Thr Ser Lys Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser
    50
                        55
                                             60
Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu
                                                             80
65
                    70
                                         75
Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Asn Pro Phe Thr
                                                         95
                85
                                     90
Phe Gly Gln Gly Thr Lys Leu Gln Ile Thr
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Arg Ala Ser Ser Ser Val Ser Tyr Met Asn
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peptide

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caacagtgga gtagtaaccc tctcact
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atcacttgca gagcaagttc aagcgtaagc tacatgaatt ggtatcagca gacaccaggg
                                                                      180
aaagccccta agagatggat ctatgacaca tccaaagtgg cttctggggt cccatcaagg
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240 ttcagtggca gtggatctgg gacagattac actttcacca tcagcagtct gcaacctgaa gatattgcaa cttactactg tcaacagtgg agtagtaacc ctctcacttt tggccagggg 300 accaagetge agateace 318 <210> 10 <211> 106 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic CD3 VL <400> 10 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly 10 15 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Ser Ser Val Ser Tyr Met 20 30 25 Asn Trp Tyr Gln Gln Thr Pro Gly Lys Ala Pro Lys Arg Trp Ile Tyr 35 40 45 Asp Thr Ser Lys Val Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser 50 55 60 Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu 80 65 70 75 Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Asn Pro Leu Thr 90 95 Phe Gly Gln Gly Thr Lys Leu Gln Ile Thr 100 105 <210> 11 <211> 357 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic CD3 VH <400> 11 caggtgcagc tggtgcagtc tgggggaggc gtggtccagc ctgggaggtc cctgagactc 60 120 tcctgtaagt cttctggata caccttcact aggtatacga tgcactgggt ccgccaggct ccagggaagg ggctggagtg gattggatac ataaatccta gccgtggtta tactaattat 180 aatcagaagg tgaaggaccg attcaccatc tccagagaca actccaagaa cacggccttt 240 ctgcaaatgg acagcctgag acccgaggac acgggtgtgt atttctgtgc gagatattat 300

gatgatcatt actgccttga ctactggggc cagggcaccc cggtcaccgt ctcctca

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Ser	Leu	Arg	Leu 20	Ser	Cys	Lys	Ser	Ser 25	Gly	Tyr	Thr	Phe	Thr 30	Arg	Tyr
Thr	Met	His 35	Trp	Val	Arg	Gln	Ala 40	Pro	Gly	Lys	Gly	Leu 45	Glu	Trp	Ile
Gly	Tyr 50	Ile	Asn	Pro	Ser	Arg 55	Gly	Tyr	Thr	Asn	Tyr 60	Asn	Gln	Lys	Val
Lys 65	Asp	Arg	Phe	Thr	Ile 70	Ser	Arg	Asp	Asn	Ser 75	Lys	Asn	Thr	Ala	Phe 80
Leu	Gln	Met	Asp	Ser 85	Leu	Arg	Pro	Glu	Asp 90	Thr	Gly	Val	Tyr	Phe 95	Cys
Ala	Arg	Tyr	Tyr 100	Asp	Asp	His	Tyr	Cys 105	Leu	Asp	Tyr	Trp	Gly 110	Gln	Gly
Thr	Pro	Val 115	Thr	Val	Ser	Ser									
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<210> 14

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<400> 14

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Ser Leu Arg Leu Ser Cys Lys Ser Ser Gly Tyr Thr Phe Thr Arg Tyr 20 25 30

Thr Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile 35 40 45

Gly Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Val 50 60

Lys Asp Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Ala Phe 70 75 80

Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys
85 90 95

Ala Arg Tyr Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly
100 105 110

Thr Pro Val Thr Val Ser Ser Val Glu Gly Gly Ser Gly Gly Ser Gly 115 120 125

Gly Ser Gly Gly Ser Gly Gly Val Asp Asp Ile Gln Met Thr Gln Ser 130 135 140

Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys 145 150 155 160 Arg Ala Ser Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Thr Pro 165 170 175 Gly Lys Ala Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser 180 185 190 Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr 195 200 205 Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys 210 215 220 Gln Gln Trp Ser Ser Asn Pro Leu Thr Phe Gly Gln Gly Thr Lys Leu 225 240 230 235 Gln Ile Thr <210> 15 <211> 372 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic CD19 VH <400> 15 60 caggtgcagc tgcagcagtc tggggctgag ctggtgaggc ctgggtcctc agtgaagatt 120 tcctgcaagg cttctggcta tgcattcagt agctactgga tgaactgggt gaagcagagg cctggacagg gtcttgagtg gattggacag atttggcctg gagatggtga tactaactac 180 aatggaaagt tcaagggtaa agccactctg actgcagacg aatcctccag cacagcctac 240 300 atgcaactca gcagcctagc atctgaggac tctgcggtct atttctgtgc aagacgggag actacgacgg taggccgtta ttactatgct atggactact ggggccaagg gaccacggtc 360 372 accgtctcct cc <210> 16 <211> 124 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic CD19 VH

<400> 16

<213> Artificial Sequence

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                                     10
Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Tyr
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                                 25
                                                     30
Trp Met Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35
                             40
                                                 45
Gly Gln Ile Trp Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe
    50
                         55
                                             60
Lys Gly Lys Ala Thr Leu Thr Ala Asp Glu Ser Ser Ser Thr Ala Tyr
                                                             80
65
                    70
                                         75
Met Gln Leu Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Phe Cys
                                     90
                85
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Ala Arg Arg Glu Thr Thr Thr Val Gly Arg Tyr Tyr Ala Met Asp
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                                 105
Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser
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                                                                      120
                                                                      180
caacagattc caggacagcc acccaaactc ctcatctatg atgcatccaa tctagtttct
gggatcccac ccaggtttag tggcagtggg tctgggacag acttcaccct caacatccat
                                                                      240
cctgtggaga aggtggatgc tgcaacctat cactgtcagc aaagtactga ggatccgtgg
                                                                      300
                                                                      333
acgttcggtg gagggaccaa gctcgagatc aaa
<210> 18
<211> 111
<212> PRT
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<220>

<223> Description of Artificial Sequence: Synthetic CD19 VL

<400> 18

Asp Ile Gln Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly 1 5 15

Gln Arg Ala Thr Ile Ser Cys Lys Ala Ser Gln Ser Val Asp Tyr Asp 20 25 30

Gly Asp Ser Tyr Leu Asn Trp Tyr Gln Gln Ile Pro Gly Gln Pro Pro 35 40 45

Lys Leu Leu Ile Tyr Asp Ala Ser Asn Leu Val Ser Gly Ile Pro Pro 50 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His 70 75 80

Pro Val Glu Lys Val Asp Ala Ala Thr Tyr His Cys Gln Gln Ser Thr 85 90 95

Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105 110

<210> 19

<211> 1504

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 anti-CD3

<400> 19

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1504

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aatggacagc ctgagacccg aggacacggg tgtgtatttc tgtgcgagat attatgatga 1080
tcattactgc cttgactatt ggggccaggg caccccggtc accgtctcct cagtcgaagg 1140
tggaagtgga ggttctggtg gaagtggagg ttcaggtgga gtggacgaca tccagatgac 1200
ccagtctcca tcctccctgt ctgcatctgt aggagacaga gtcaccatca cttgcagagc 1260
aagttcaagc gtaagctaca tgaattggta tcagcagaca ccagggaaag cccctaagag 1320
atggatctat gacacatcca aagtggcttc tggggtccca tcaaggttca gtggcagtgg 1380
atctgggaca gattacactt tcaccatcag cagtctgcaa cctgaagata ttgcaactta 1440
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<210> 20
<211> 498
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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                                    10
Gln Arg Ala Thr Ile Ser Cys Lys Ala Ser Gln Ser Val Asp Tyr Asp
            20
                                25
                                                     30
Gly Asp Ser Tyr Leu Asn Trp Tyr Gln Gln Ile Pro Gly Gln Pro Pro
                                                 45
        35
                            40
Lys Leu Leu Ile Tyr Asp Ala Ser Asn Leu Val Ser Gly Ile Pro Pro
    50
                        55
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His
                                                             80
65
                    70
                                        75
Pro Val Glu Lys Val Asp Ala Ala Thr Tyr His Cys Gln Gln Ser Thr
                85
                                    90
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Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Gly 105 110 100

Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Gln Val 125 115 120

Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ser Ser Val 135 130 140

Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Tyr Trp Met
145 150 155 160

Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly Gln 175

Ile Trp Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys Gly 180 185 190

Lys Ala Thr Leu Thr Ala Asp Glu Ser Ser Ser Thr Ala Tyr Met Gln
195 200 205

Leu Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Phe Cys Ala Arg 210 215 220

Arg Glu Thr Thr Val Gly Arg Tyr Tyr Ala Met Asp Tyr Trp 225 230 235 240

Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gln 245 250 255

Val Gln Leu Val Gln Ser Gly Gly Gly Val Val Gln Pro Gly Arg Ser 260 265 270

Leu Arg Leu Ser Cys Lys Ser Ser Gly Tyr Thr Phe Thr Arg Tyr Thr 275 280 285

Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly 290 295 300

Tyr Ile Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Val Lys 305 310 315 320

Asp Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Ala Phe Leu 325 330 335

Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys Ala 340 345 350

Arg Tyr Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly Thr 355 360 365

Pro Val Thr Val Ser Ser Val Glu Gly Gly Ser Gly Gly Ser Gly Gly 370 380

Ser Gly Gly Ser Gly Gly Val Asp Asp Ile Gln Met Thr Gln Ser Pro 385 390 395 400

Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg 405 410 415

Ala Ser Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Thr Pro Gly 420 425 430

Lys Ala Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser Gly 435 440 445

Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe 450 455 460

Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln 465 470 475 480

Gln Trp Ser Ser Asn Pro Leu Thr Phe Gly Gln Gly Thr Lys Leu Gln 485 490 495

Ile Thr

<210> 21

<211> 360

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic 5-10 VH

<400> 21

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<210> 22

<211> 120

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic 5-10 VH <400> 22 Glu Val Gln Leu Leu Glu Gln Ser Gly Ala Glu Leu Val Arg Pro Gly 5 10 15 Thr Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Thr Asn 20 25 30 Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu Trp 35 40 45 Ile Gly Asp Ile Phe Pro Gly Ser Gly Asn Ile His Tyr Asn Glu Lys 50 55 Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala 80 65 70 75 Tyr Met Gln Leu Ser Ser Leu Thr Phe Glu Asp Ser Ala Val Tyr Phe 95 85 90 Cys Ala Arg Leu Arg Asn Trp Asp Glu Pro Met Asp Tyr Trp Gly Gln 100 110 105 Gly Thr Thr Val Thr Val Ser Ser 115 120 <210> 23 <211> 339 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 5-10 VL <400> 23 60 gagetegtga tgacacagte tecateetee etgactgtga cageaggaga gaaggteaet 120 atgagetgea agtecagtea gagtetgtta aacagtggaa atcaaaagaa etaettgace 180 tggtaccagc agaaaccagg gcagcctcct aaactgttga tctactgggc atccactagg gaatctgggg tccctgatcg cttcacaggc agtggatctg gaacagattt cactctcacc 240 atcagcagtg tgcaggctga agacctggca gtttattact gtcagaatga ttatagttat 300 ccgctcacgt tcggtgctgg gaccaagctt gagatcaaa 339

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<210> 24
<211> 113
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Asn Ser
            20
                                 25
                                                     30
Gly Asn Gln Lys Asn Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln
        35
                                                 45
                             40
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
    50
                         55
                                             60
Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
                                                             80
                                         75
65
                    70
Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Asn
                                                         95
                                     90
                85
Asp Tyr Ser Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Ile
            100
                                 105
                                                     110
Lys
<210> 25
<211> 360
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      3-1 VH
<400> 25
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atatcctgca aggcttctgg atacgccttc actaactact ggctaggttg ggtaaagcag
                                                                       120
                                                                       180
aggcctggac atggacttga gtggattgga gatcttttcc ctggaagtgg taatactcac
                                                                       240
tacaatgaga ggttcagggg caaagccaca ctgactgcag acaaatcctc gagcacagcc
tttatgcagc tcagtagcct gacatctgag gactctgctg tctatttctg tgcaagattg
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aggaactggg acgaggctat ggactactgg ggccaaggga ccacggtcac cgtctcctcc

<210> 26

<211> 120 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 3-1 VH <400> 26 Glu Val Gln Leu Leu Glu Gln Ser Gly Ala Glu Leu Val Lys Pro Gly 5 10 1 15 Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Thr Asn 20 25 30 Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu Trp 35 40 45 Ile Gly Asp Leu Phe Pro Gly Ser Gly Asn Thr His Tyr Asn Glu Arg 50 55 Phe Arg Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala 70 80 65 75 Phe Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe 85 95 90 Cys Ala Arg Leu Arg Asn Trp Asp Glu Ala Met Asp Tyr Trp Gly Gln 100 105 110 Gly Thr Thr Val Thr Val Ser Ser 115 120 <210> 27 <211> 321 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 3-1 VL <400> 27 gagctcgtca tgacccagtc tccatcttat cttgctgcat ctcctggaga aaccattact attaattgca gggcaagtaa gagcattagc aaatatttag cctggtatca agagaaacct gggaaaacta ataagcttct tatctactct ggatccactt tgcaatctgg aattccatca aggttcagtg gcagtggatc tggtacagat ttcactctca ccatcagtag cctggagcct

60

120

180

300

gaagattttg caatgtatta ctgtcaacag cataatgaat atccgtacac gttcggaggg 321 gggaccaagc ttgagatcaa a <210> 28 <211> 107 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 3-1 VL <400> 28 Glu Leu Val Met Thr Gln Ser Pro Ser Tyr Leu Ala Ala Ser Pro Gly 10 15 1 Glu Thr Ile Thr Ile Asn Cys Arg Ala Ser Lys Ser Ile Ser Lys Tyr 20. 25 30 Leu Ala Trp Tyr Gln Glu Lys Pro Gly Lys Thr Asn Lys Leu Leu Ile 35 40 Tyr Ser Gly Ser Thr Leu Gln Ser Gly Ile Pro Ser Arg Phe Ser Gly 50 55 60 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro 75 80 65 70 Glu Asp Phe Ala Met Tyr Tyr Cys Gln Gln His Asn Glu Tyr Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys 105 100 <210> 29 <211> 372 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 4-7 VH <400> 29 gaggtgcagc tgctcgagca gtctggagct gagctggcga ggcctggggc ttcagtgaag 60 ctgtcctgca aggcttctgg ctacaccttc acaaactatg gtttaagctg ggtgaagcag 120 180 aggcctggac aggtccttga gtggattgga gaggtttatc ctagaattgg taatgcttac tacaatgaga agttcaaggg caaggccaca ctgactgcag acaaatcctc cagcacagcg 240 300 tccatggagc tccgcagcct gacctctgag gactctgcgg tctatttctg tgcaagacgg 360 ggatcctacg atactaacta cgactggtac ttcgatgtct ggggccaagg gaccacggtc

372 accgtctcct cc <210> 30 <211> 124 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 4-7 VH <400> 30 Glu Val Gln Leu Leu Glu Gln Ser Gly Ala Glu Leu Ala Arg Pro Gly 5 10 15 1 Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asn 30 25 20 Tyr Gly Leu Ser Trp Val Lys Gln Arg Pro Gly Gln Val Leu Glu Trp 40 45 35 Ile Gly Glu Val Tyr Pro Arg Ile Gly Asn Ala Tyr Tyr Asn Glu Lys 60 50 55 Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala 80 70 75 65 Ser Met Glu Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe 95 90 85 Cys Ala Arg Arg Gly Ser Tyr Asp Thr Asn Tyr Asp Trp Tyr Phe Asp 105 110 100 Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser 115 120 <210> 31 <211> 336 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 4-7 VL <400> 31 gagetegtga tgacceagae tecaetetee etgeetgtea gtettggaga teaageetee 60 atctcttgca gatctagtca gagccttgta cacagtaatg gaaacaccta tttacattgg 120 tacctgcaga agccaggcca gtctccaaag ctcctgatct acaaagtttc caaccgattt 180

240 totggggtcc cagacaggtt cagtggcagt ggatcaggga cagatttcac actcaagatc agcagagtgg aggctgagga tctgggagtt tatttctgct ctcaaagtac acatgttccg 300 tacacgttcg gagggggac caagcttgag atcaaa 336 <210> 32 <211> 112 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic 4-7 VL <400> 32 Glu Leu Val Met Thr Gln Thr Pro Leu Ser Leu Pro Val Ser Leu Gly 15 10 1 Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser 25 30 20 Asn Gly Asn Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser 35 40 45 Pro Lys Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro 50 55 60 Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile 80 65 70 75 Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser 85 90 Thr His Val Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys 105 110 100 <210> 33 <211> 1470 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic anti-CD3 <400> 33 gagetegtea tgacceagte tecatettat ettgetgeat etectggaga aaceattaet 60 attaattgca gggcaagtaa gagcattagc aaatatttag cctggtatca agagaaacct 120 gggaaaacta ataagcttct tatctactct ggatccactt tgcaatctgg aattccatca 180 aggttcagtg gcagtggatc tggtacagat ttcactctca ccatcagtag cctggagcct 240 gaagattttg caatgtatta ctgtcaacag cataatgaat atccgtacac gttcggaggg 300 gggaccaage ttgagatcaa aggtggtggt ggttctggcg gcggcggctc cggtggtggt ggttctgagg tgcagctgct cgagcagtct ggagctgagc tggtgaaacc tggggcctca

360

420

480

540

600

660

720

780

840

900

960

1020

1080

1140

1200

1260

1320

1380

1440

1470

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gtgaagatat cctgcaaggc ttctggatac gccttcacta actactggct aggttgggta
aagcagaggc ctggacatgg acttgagtgg attggagatc ttttccctgg aagtggtaat
actcactaca atgagaggtt caggggcaaa gccacactga ctgcagacaa atcctcgagc
acageettta tgeageteag tageetgaea tetgaggaet etgetgteta tttetgtgea
agattgagga actgggacga ggctatggac tactggggcc aagggaccac ggtcaccgtc
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tatttctgtg cgagatatta tgatgatcat tactgccttg actattgggg ccagggcacc
ccggtcaccg tctcctcagt cgaaggtgga agtggaggtt ctggtggaag tggaggttca
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gtcccatcaa ggttcagtgg cagtggatct gggacagatt acactttcac catcagcagt
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<210> 34
<211> 490
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      anti-CD3
<400> 34
Glu Leu Val Met Thr Gln Ser Pro Ser Tyr Leu Ala Ala Ser Pro Gly
                                    10
                                                         15
                5
1
Glu Thr Ile Thr Ile Asn Cys Arg Ala Ser Lys Ser Ile Ser Lys Tyr
            20
                                25
Leu Ala Trp Tyr Gln Glu Lys Pro Gly Lys Thr Asn Lys Leu Leu Ile
                                                 45
        35
Tyr Ser Gly Ser Thr Leu Gln Ser Gly Ile Pro Ser Arg Phe Ser Gly
                                             60
    50
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
                                                             80
                                         75
                    70
65
Glu Asp Phe Ala Met Tyr Tyr Cys Gln Gln His Asn Glu Tyr Pro Tyr
                                                         95
                                     90
                85
Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Gly Gly Gly Ser
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105

100

Gly Gly Gly Ser Gly Gly Gly Ser Glu Val Gln Leu Leu Glu 115 120 125

Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala Ser Val Lys Ile Ser 130 135 140

Cys Lys Ala Ser Gly Tyr Ala Phe Thr Asn Tyr Trp Leu Gly Trp Val 145 150 150

Lys Gln Arg Pro Gly His Gly Leu Glu Trp Ile Gly Asp Leu Phe Pro 165 170 175

Gly Ser Gly Asn Thr His Tyr Asn Glu Arg Phe Arg Gly Lys Ala Thr 180 185 190

Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Phe Met Gln Leu Ser Ser 195 200 205

Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys Ala Arg Leu Arg Asn 210 215 220

Trp Asp Glu Ala Met Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val 225 230 235 240

Ser Ser Gly Gly Gly Ser Gln Val Gln Leu Val Gln Ser Gly Gly 245 250 255

Gly Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Lys Ser Ser 260 265 270

Gly Tyr Thr Phe Thr Arg Tyr Thr Met His Trp Val Arg Gln Ala Pro 275 280 285

Gly Lys Gly Leu Glu Trp Ile Gly Tyr Ile Asn Pro Ser Arg Gly Tyr 290 295 300

Thr Asn Tyr Asn Gln Lys Val Lys Asp Arg Phe Thr Ile Ser Arg Asp 305 310 315 320

Asn Ser Lys Asn Thr Ala Phe Leu Gln Met Asp Ser Leu Arg Pro Glu 325 330 335

Asp Thr Gly Val Tyr Phe Cys Ala Arg Tyr Tyr Asp Asp His Tyr Cys 340 345 350

Leu Asp Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Val Glu 355 360 365

Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Gly Val Asp 370 375 380

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly 385 390 395 400

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Ser Ser Val Ser Tyr Met 405 410 415

Asn Trp Tyr Gln Gln Thr Pro Gly Lys Ala Pro Lys Arg Trp Ile Tyr 420 425 430

Asp Thr Ser Lys Val Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser 435 440 445

Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu 450 455 460

Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Asn Pro Leu Thr 465 470 475 480

Phe Gly Gln Gly Thr Lys Leu Gln Ile Thr 485

<210> 35

<211> 1498

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic anti-CD3

<400> 35

tgtacactcc gagctcgtga tgacacagtc tccatcctcc ctgactgtga cagcaggaga 60 gaaggtcact atgagctgca agtccagtca gagtctgtta aacagtggaa atcaaaagaa 120 ctacttgacc tggtaccagc agaaaccagg gcagcctcct aaactgttga tctactgggc 180 atccactagg gaatctgggg tccctgatcg cttcacaggc agtggatctg gaacagattt 240 cactctcacc atcagcagtg tgcaggctga agacctggca gtttattact gtcagaatga 300 ttatagttat ccgctcacgt tcggtgctgg gaccaagctt gagatcaaag gtggtggtgg 360 ttctggcggc ggcggctccg gtggtggtgg ttctgaggtg cagctgctcg agcagtctgg 420 agctgagctg gtaaggcctg ggacttcagt gagatatcc tgcaaggctt ctggatacgc 480

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cttcactaac tactggctag gttgggtaaa gcagaggcct ggacatggac ttgagtggat 540
tggagatatt ttccctggaa gtggtaatat ccactacaat gagaagttca agggcaaagc 600
cacactgact gcagacaaat cttcgagcac agcctatatg cagctcagta gcctgacatt 660
tgaggactct gctgtctatt tctgtgcaag actgaggaac tgggacgagc ctatggacta 720
ctggggccaa gggaccacgg tcaccgtctc ctccggaggt ggtggctccc aggtgcagct 780
ggtgcagtct gggggaggcg tggtccagcc tgggaggtcc ctgagactct cctgtaagtc 840
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gctggagtgg attggataca taaatcctag ccgtggttat actaattata atcagaaggt 960
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tggaggttct ggtggaagtg gaggttcagg tggagtggac gacatccaga tgacccagtc 1200
tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc atcacttgca gagcaagttc 1260
aagcgtaagc tacatgaatt ggtatcagca gacaccaggg aaagccccta agagatggat 1320
ctatgacaca tccaaagtgg cttctggggt cccatcaagg ttcagtggca gtggatctgg 1380
gacagattac actttcacca tcagcagtct gcaacctgaa gatattgcaa cttactactg 1440
tcaacagtgg agtagtaacc ctctcacttt tggccagggg accaagctgc agatcacc
                                                                  1498
<210> 36
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<21U> 36
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<211> 496

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic anti-CD3

<400> 36

Glu Leu Val Met Thr Gln Ser Pro Ser Ser Leu Thr Val Thr Ala Gly
1 5 10

Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Asn Ser 20 25 30

Gly Asn Gln Lys Asn Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln 35 40 45

Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val 50 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr 65 70 75 80

Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Asn 85 90 95

Asp Tyr Ser Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Ile 100 105 110 Lys Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser 115 120 125

Glu Val Gln Leu Leu Glu Gln Ser Gly Ala Glu Leu Val Arg Pro Gly 130 135 140

Thr Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Thr Asn 145 150 155 160

Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu Trp
165 170 175

Ile Gly Asp Ile Phe Pro Gly Ser Gly Asn Ile His Tyr Asn Glu Lys
180 185 190

Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala 195 200 205

Tyr Met Gln Leu Ser Ser Leu Thr Phe Glu Asp Ser Ala Val Tyr Phe 210 220

Cys Ala Arg Leu Arg Asn Trp Asp Glu Pro Met Asp Tyr Trp Gly Gln 225 230 235 240

Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gln Val Gln 245. 250 255

Leu Val Gln Ser Gly Gly Gly Val Val Gln Pro Gly Arg Ser Leu Arg 260 265 270

Leu Ser Cys Lys Ser Ser Gly Tyr Thr Phe Thr Arg Tyr Thr Met His 275 280 285

Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly Tyr Ile 290 295 300

Asn Pro Ser Arg Gly Tyr Thr Asn Tyr Asn Gln Lys Val Lys Asp Arg 305 310 315 320

Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Ala Phe Leu Gln Met 325 330 335

Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys Ala Arg Tyr 340 345 350

Tyr Asp Asp His Tyr Cys Leu Asp Tyr Trp Gly Gln Gly Thr Pro Val 355 360 365

Thr Val Ser Ser Val Glu Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly 370 380

Gly Ser Gly Gly Val Asp Asp Ile Gln Met Thr Gln Ser Pro Ser Ser 385 390 395 400

Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser 405 410 415

Ser Ser Val Ser Tyr Met Asn Trp Tyr Gln Gln Thr Pro Gly Lys Ala 420 425 430

Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Val Ala Ser Gly Val Pro 435 440 445

Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr Phe Thr Ile 450 455 460

Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Trp 465 470 475 480

Ser Ser Asn Pro Leu Thr Phe Gly Gln Gly Thr Lys Leu Gln Ile Thr 485 490 495

<210> 37

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 37

Leu Glu Trp Ile Gly
1 5

<210> 38

<211> 10

<212> PRT

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 38
Ser Ala Ser Ser Ser Val Ser Tyr Met Asn
                                     10
  1
<210> 39
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 39
Asp Thr Ser Lys Leu Ala Ser
  1
<210> 40
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 40
Gln Gln Trp Ser Ser Asn Pro Phe Thr
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5